

Amendments to the Specification:

Table 2: Oligonucleotides for detecting the expression of human chemokine receptors

Receptor	EMBL nucleotide sequence database identifier	Amplificate length (base pairs)	Oligonucleotide sequence (5'>3')
ccr1 upper	NM 001295	129	GAGCCAATCAGTAGGCCAGCATCT (SEQ ID NO. 1)
ccr1 lower	NM 001295		GTTCCCCATTTCTATTTCTCGTT (SEQ ID NO. 2)
ccr2 upper	NM 000647	173	CTCCCTGAAGTAAGCAAAGAC (SEQ ID NO. 3)
ccr2 lower	NM 000647		CCATGTGGCCTGAAAGTAG (SEQ ID NO. 4)
ccr3 upper	NM 178329	148	GGCAGATACATCCATTCCCTTC (SEQ ID NO. 5)
ccr3 lower	NM 178329		GGTTGCTTCATCTCCTTGGTCCTT (SEQ ID NO. 6)
ccr4 upper	X85740	91	CAGGGGCCTTTGTGCTC (SEQ ID NO. 7)
ccr4 lower	X85740		CATGGTGGACTGCGTGTAAGAT (SEQ ID NO. 8)
ccr5 upper	NM 000579	160	AGGAGGGAGGTATTCGTAAGG (SEQ ID NO. 9)
ccr5 lower	NM 000579		TTCAAGGGTTCTCCAATCTG (SEQ ID NO. 10)
ccr6 upper	NM 031409	86	TGGTTACAGCACAAAATGATGG (SEQ ID NO. 11)
ccr6 lower	NM 031409		TTGCCTAAATGAGTGATGTGTTG (SEQ ID NO. 12)
ccr7 upper	NM 001838	194	GCCGCCCTAAAAGCACACTCATCC (SEQ ID NO. 13)
ccr7 lower	NM 001838		TTCCCTTGCCTCTCCTCCCATCC (SEQ ID NO. 14)
ccr8 upper	NM 005201	198	TGCAGCCAAATCTCAACTACC (SEQ ID NO. 15)
ccr8 lower	NM 005201		AAACCTTCACACCCACACCTT (SEQ ID NO. 16)
ccr9 upper	NM 031200	151	AGCCTTGGCCCTGTTGTA (SEQ ID NO. 17)
ccr9 lower	NM 031200		TGCCCATATCTGCTCACTGTA (SEQ ID NO. 18)
ccr10 upper	NM 016602	118	GCCCCGCCCTTCTTCCTGCTCA (SEQ ID NO. 19)
ccr10 lower	NM 016602		CCACCTACTCCCCTTCCCACGAC (SEQ ID NO. 20)
ccr11 upper	NM 016557	90	CTCTGCCTTTGCTTGGATACATA (SEQ ID NO. 21)
ccr11 lower	NM 016557		CACGGCGTCTGAGATTGAGTT (SEQ ID NO. 22)
cxcr1 upper	NM 000634	177	CCGTGCTTGTCCCTGTGG (SEQ ID NO. 23)
cxcr1 lower	NM 000634		CTGTGCCTCAAGAGACTGTT (SEQ ID NO. 24)
cxcr2 upper	NM 001557	146	AGTTTATGATTCCACCTACA (SEQ ID NO. 25)

cxcr2 lower	NM 001557		TTCAACATCCTAACATAAA (<u>SEQ ID NO. 26</u>)
cxcr3 upper	NM 001504	140	GTGGCCGAGAAAGCAGGGTAGACG (<u>SEQ ID NO. 27</u>)
cxcr3 lower	NM 001504		CAGGCGCAAGAGCAGCATCCACAT (<u>SEQ ID NO. 28</u>)
cxcr4 upper	NM 003467	141	GATCCCTGCCCTCCTGCTGACTAT (<u>SEQ ID NO. 29</u>)
cxcr4 lower	NM 003467		AGGCCAACCATGATGTGCTGAAAC (<u>SEQ ID NO. 30</u>)
cxcr5 upper	NM 032966	170	CCGGATCCTGGGTGGTCTG (<u>SEQ ID NO. 31</u>)
cxcr5 lower	NM 032966		CCGCCGGTTTGATTGAT (<u>SEQ ID NO. 32</u>)
cxcr6 upper	NM 006564	119	GACTTTCCCTCCTCCATCTCCA (<u>SEQ ID NO. 33</u>)
cxcr6 lower	NM 006564		GGCCGTGCTCACCTCTCA (<u>SEQ ID NO. 34</u>)
Cx3cr upper	NM 001337	169	TAGGCCAAGTTGTATCAGGTG (<u>SEQ ID NO. 35</u>)
Cx3cr lower	NM 001337		GTGTGGCATTGTTGTGTA (<u>SEQ ID NO. 36</u>)
xcr upper	NM 005283	181	AGCTCATCTCGCCATCGTG (<u>SEQ ID NO. 37</u>)
xcr lower	NM 005283		ACCGGGTTAAAGCAGCAGTG (<u>SEQ ID NO. 38</u>)